

REMARKS

This supplemental response was requested by the examiner in a telephone conference on May 21, 2010. As requested, Applicants have amended claims 67, 73, 74, 255, and 278 to clarify the claim language. In this regard, Applicants note that the claims presented herein are amended relative to the claim amendments entered in the response of March 8, 2010. In addition, Applicants have considered the similarity of certain amino acid sequences for the HPIV2 L protein provided by the examiner and present the results of the assessment below. No new matter is added in this response.

Assessment of Amino Acid Sequences Reported for the HPIV2 L Protein:

During the conference call of May 21st, the examiner expressed concern that other HIPV2 protein sequences (GenBank accession numbers NP_598406.1, CAA40788.1, and BAE00056.1) might not have sufficient homology to the sequences disclosed in the application to warrant the scope of the claims. Applicants contend that any such concerns are misplaced.

The sequences corresponding to the accession numbers above were compared via sequence alignment using the default settings of the SIM program of the Expert Protein Analysis System (ExPASy) – available at <http://expasy.org/tools/sim-prot.html>. The sequences for accession numbers NP_598406.1 and CAA40788.1 were found to be 100% identical over their entire 2262 amino acid sequences (the results of this alignment are provided on pages 6-9). Similarly, sequences for accession number BAE00056.1 was found to be 99.3% identical to NP_598406.1 and CAA40788.1, with 15 sequence differences in 2264 amino acids (the results of this alignment are provided on pages 10-13; however, because of the identity of NP_598406.1 and CAA40788.1, the alignment only compares NP_598406.1 with BAE00056.1). Nonetheless, the relevant segments of the sequences for accession numbers NP_598406.1, CAA40788.1, and BAE00056.1 are 100% identical to the HIPV2 L protein amino acid sequences shown in figure 1 of the application.

In comparing the HPIV2 L protein sequences, Applicants noticed that the amino acid numbering of the NP_598406.1 and CAA40788.1 sequences differs slightly from that of BAE00056.1 and figure 1. In particular, residues 948 and 1566 of figure 1 are numbered as

residues 946 and 1564, respectively, likely due to the presence of residue insertions in the sequence disclosed in the present application. Therefore, Applicants have amended claims 67, 255, and 278 to clarify the scope of the claims. In this regard, the claims now recite residues of the SEQ ID NOs, that correspond to the relevant sequences of figure 1. Applicants note that the claimed SEQ IN NOs, were submitted for the pending application on May 13, 2004. Support for the amended claims can be found throughout the specification, such as in paragraph [39] and figure 1.


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Results of SIMsearch:

Sequence 1: NP_598406.1 (2757 residues)
 Sequence 2: CA440789.1 (2757 residues)

Using the parameters:

Comparison matrix: BLOSUM62
 Minimum of alignment score: 20
 Gap open penalty: 12
 Gap extension penalty: 4



Evaluate the significance of this protein sequence similarity score using PRSS at EMBL-CH.

100.0% identity in 2262 residues overlap; Score: 11757.0; Gap frequency: 0.0%

NP_598406.1	1	MAASSTILLPEVHLNPFVHKLITYLLGHFPHOLDIGHSISLHNDWQIIMSESRLE
CA440789.1	1	MAASSTILLPEVHLNPFVHKLITYLLGHFPHOLDIGHSISLHNDWQIIMSESRLE

NP_598406.1	61	ERLOVAKZELIKYVPAPFATWBSHAWLIWPSCTPELKKPLPKLQWSONYKLINAS
CA440789.1	61	ERLOVAKZELIKYVPAPFATWBSHAWLIWPSCTPELKKPLPKLQWSONYKLINAS

NP_598406.1	121	CNTTSLGQTRCMENISITLGTGHELPSSRGTWACWASKITLMDTSINSESHKQSPUS
CA440789.1	121	CNTTSLGQTRCMENISITLGTGHELPSSRGTWACWASKITLMDTSINSESHKQSPUS

NP_598406.1	181	EWLTIKYPOMQLNHQSGROFTLVHIVTNSQLVITPELVICPDFLNSVLMYPTPTNT
CA440789.1	181	EWLTIKYPOMQLNHQSGROFTLVHIVTNSQLVITPELVICPDFLNSVLMYPTPTNT

NP_598406.1	241	LNPSTMPFGEHMYTILCTISHVLSVLGPRIDALFSTVDLALGLQIVYKVIASLESLEY
CA440789.1	241	LNPSTMPFGEHMYTILCTISHVLSVLGPRIDALFSTVDLALGLQIVYKVIASLESLEY

NP-598406. 361 GCGGLADPWVTEAGSFRSFTQETIIDILIGSKGLDFRSTIYVTTQALLIPSHUSPLALAE
CAA49789.1 362 GCGGLADPWVTEAGSFRSFTQETIIDILIGSKGLDFRSTIYVTTQALLIPSHUSPLALAE

NP-598406 381 MLCGMRGNGHPTLAAQVAVNRKSHCAKFLLEPFTKTKLFPFHTILIDGTRKKKMGHAF
CAA49789.1 382 MLCGMRGNGHPTLAAQVAVNRKSHCAKFLLEPFTKTKLFPFHTILIDGTRKKKMGHAF

NP-598406. 401 ELILFVRASYSILIESQMDRAKISYEYTLKMKKILSLIEPRCCTFDFGNEELSIPIINDMAI
CAA49789.1 402 ELILFVRASYSILIESQMDRAKISYEYTLKMKKILSLIEPRCCTFDFGNEELSIPIINDMAI

NP-598406. 481 SAPREDWMEVFKKSLIAQPRGPHIPHPHPMKRLLLNPLSDSDFCPVAKLRYVIGWEYL
CAA49789.1 482 SAPREDWMEVFKKSLIAQPRGPHIPHPHPMKRLLLNPLSDSDFCPVAKLRYVIGWEYL

NP-598406. 541 QDDTFCASYSLEKEIKFDGRIFAKLITRMRSQGVIAKAILAHNAGTLMKKNIVVLNQLS
CAA49789.1 542 QDDTFCASYSLEKEIKFDGRIFAKLITRMRSQGVIAKAILAHNAGTLMKKNIVVLNQLS

NP-598406. 601 LTKGLITMIGIGIILSEKAKFTYTDNHSQGVFTYITWDSKMKRKSCTAGSVLADPDVFPAL
CAA49789.1 602 LTKGLITMIGIGIILSEKAKFTYTDNHSQGVFTYITWDSKMKRKSCTAGSVLADPDVFPAL

NP-598406 681 SACTYITIDLAKYCYLQWRYGTLIRFARTLHMVYVFLFESVILGLIRSTLYVDFDFPFA
CAA49789.1 682 SACTYITIDLAKYCYLQWRYGTLIRFARTLHMVYVFLFESVILGLIRSTLYVDFDFPFA

NP-598406. 721 AIDAVDLQKVLGGDPIFVAGSGIEELQGMWNTIPLSVLILGEAGETVRYSTMYGKRNQA
CAA49789.1 722 AIDAVDLQKVLGGDPIFVAGSGIEELQGMWNTIPLSVLILGEAGETVRYSTMYGKRNQA

NP-598406. 781 IAVTTPVPRKLPSIQKKELAYAAKSLFFRELRAHMYGLQGLYAGETILISSTFFIYSPV
CAA49789.1 782 IAVTTPVPRKLPSIQKKELAYAAKSLFFRELRAHMYGLQGLYAGETILISSTFFIYSPV

NP-598406. 841 FYQGRILTQALKKASKLCTADVLGECTJASCSNSATTINRLTENGVERDTCYKLNITYGS
CAA49789.1 842 FYQGRILTQALKKASKLCTADVLGECTJASCSNSATTINRLTENGVERDTCYKLNITYGS

NP-598406. 901 INQLTYLLIFFQYSIPKCTISSIFLQPKLISKILVILPSQLGGSLMYLACSKLNNKICA
CAA49789.1 902 INQLTYLLIFFQYSIPKCTISSIFLQPKLISKILVILPSQLGGSLMYLACSKLNNKICA

NP-598406. 961 LGTAVARLRLKNGALRESNLYNLLARFPKSGSWETLAADVYLSGQYLYFPPTILKAK
CAA49789.1 962 LGTAVARLRLKNGALRESNLYNLLARFPKSGSWETLAADVYLSGQYLYFPPTILKAK

NP-598406. 1021 TQNTLARIORFNLKSVPTNAREKEEMILLARPLADIVLPVAVIILIQSGISGNKQIQ
CAA49789.1 1022 TQNTLARIORFNLKSVPTNAREKEEMILLARPLADIVLPVAVIILIQSGISGNKQIQ

NP-598406. 1081 GNPFTIPTIMPRESFKLELTKKYLIVKRYGTYLNVNVPVILNLELFPYSLNLYITIQTC

CAA40708.1 1501 GFFDTTPTIMRHSPEIKPLSTKPLSLVSYENUTLSYVYFVILALPLFLPGLYIM LDUQC
 NF-598406 1141 SIDSRSLEKLSWGLLNKRLSLKTEDEFIKVVPFPLVGRDODCPMGDAKPTWFFL
 CAA40708.1 1141 SIDSRSLEKLSWGLLNKRLSLKTEDEFIKVVPFPLVGRDODCPMGDAKPTWFFL
 NF-598406 1201 RWSIILDGUPETNPPPIKVPYVIGSRTREGRVASHAYIKGATHSLKALRGAGVYLKAFQDT
 CAA40708.1 1201 RWSIILDGUPETNPPPIKVPYVIGSRTREGRVASHAYIKGATHSLKALRGAGVYLKAFQDT
 NF-598406 1261 VVMNGLDIDIANTRYKISLEQLQLTFLPLTSANITHELDGATLEKTPASSVAFSSYTH
 CAA40708.1 1261 VVMNGLDIDIANTRYKISLEQLQLTFLPLTSANITHELDGATLEKTPASSVAFSSYTH
 NF-598406 1321 LSEMQYLYELIQYVVLGHLIYQQLNITGLHIRTTHNPPISTTLEITLHLATSSCCVR
 CAA40708.1 1321 LSEMQYLYELIQYVVLGHLIYQQLNITGLHIRTTHNPPISTTLEITLHLATSSCCVR
 NF-598406 1381 SVDGCLICESNGEVFQITVPTPTTFVYDPMPLADYEIARLDNLSYQAKISTOYVYLTOR
 CAA40708.1 1381 SVDGCLICESNGEVFQITVPTPTTFVYDPMPLADYEIARLDNLSYQAKISTOYVYLTOR
 NF-598406 1441 IDLLAHLTAQKMINSTIGLDETYSIVMDAVILSDYTHWITSECSYKIELVFKLMAHNL
 CAA40708.1 1441 IDLLAHLTAQKMINSTIGLDETYSIVMDAVILSDYTHWITSECSYKIELVFKLMAHNL
 UP-598406 1501 LELAFQNYVLIISQWNTIPDYTYMTERIPSTALNNIATISHPKLLKRAHGLDITTPIN
 CAA40708.1 1501 LELAFQNYVLIISQWNTIPDYTYMTERIPSTALNNIATISHPKLLKRAHGLDITTPIN
 NF-598406 1561 APTLASLDYVQLSILSIQNGVEQVLDLSNGIDLEILLGDSMETIDPAMHLIARKLTL
 CAA40708.1 1561 APTLASLDYVQLSILSIQNGVEQVLDLSNGIDLEILLGDSMETIDPAMHLIARKLTL
 UP-598406 1621 LALYKGEHTYTFPKLKGVPFEKCLVLIEYLANCYGNTHHLDPLQKLYLNLIMKLTAF
 CAA40708.1 1621 LALYKGEHTYTFPKLKGVPFEKCLVLIEYLANCYGNTHHLDPLQKLYLNLIMKLTAF
 NF-598406 1681 SNPFYLTREILNQIKESLDEQYIITSYYESFEQLETDILHGLITAPYHNSGNHVRPL
 CAA40708.1 1681 SNPFYLTREILNQIKESLDEQYIITSYYESFEQLETDILHGLITAPYHNSGNHVRPL
 NF-598406 1741 PFDLAFHPESELEFPPLVDMHQAISATSLIOPDSNVHPLQVASTAWYKGLSYRYLLE
 CAA40708.1 1741 PFDLAFHPESELEFPPLVDMHQAISATSLIOPDSNVHPLQVASTAWYKGLSYRYLLE
 NF-598406 1801 TQKIQTDRLNLAWGSGASMSLLLELFFGLTVYINSLFESSEHPFQRNYAFLEPTQVQGV
 CAA40708.1 1801 TQKIQTDRLNLAWGSGASMSLLLELFFGLTVYINSLFESSEHPFQRNYAFLEPTQVQGV
 NF-598406 1861 PYKLNQADLADQMLIDPVPLMRGNQAVTDLTKGSAVAFIHKVBAEKALVHIDLEST
 CAA40708.1 1861 PYKLNQADLADQMLIDPVPLMRGNQAVTDLTKGSAVAFIHKVBAEKALVHIDLEST

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*****
NP-598406. 1921 ARINQQTLSRQQLNSLIATTVLKRGNILYNTSWLFFSRPSQLAGLLWQFPTRIHLIRS
CAA40788.1 1921 ANINQQTLSRQQTESLIATTVLKRFGGILIYKTEWLPSP.FGQASGLIACFFDRIMLIRS
*****

NP-598406. 1951 SYSDPFSRREVVLVCLAADEFETLRFSAALVTATTLNHGGFTTIRPTVVCSTWQRLNENVG
CAA40788.1 1951 SYSDPFSRREVVLVCLAADEFETLRFSAALVTATTLNHGGFTTIRPTVVCSTWQRLNENVG
*****

NP-598406. 2041 RVSEVILDEILDGLATNFFAGDNGILRCGGTFPSRFWLEIDQLASTDVLQDALUTLITIN
CAA40788.1 2041 RVSEVILDEILDGLATNFFAGDNGILRCGGTFPSRFWLEIDQLASTDVLQDALUTLITIN
*****

NP-598406. 2101 LKEETLVQGSNTEDYTSLLPTFTYNIGAGKVFRTIILKILERSLNVTVRNVLVLPSSIRUS
CAA40788.1 2101 LKEETLVQGSNTEDYTSLLPTFTYNIGAGKVFRTIILKILERSLNVTVRNVLVLPSSIRUS
*****

NP-598406. 2151 VRQDLSELGSRFLMSILSEQTFLEKKTFTNGVLLDQLTRIVISTFPMCHSVLFLMRPYQXLI
CAA40788.1 2151 VRQDLSELGSRFLMSILSEQTFLEKKTFTKRYLLOQLTFTYISTFPMCHSVLFLMRPYQXLI
*****

NP-598406. 2221 WKALGSVIVCSETVDIPLIKDIQIEINDFEELERGIQDEEL
CAA40788.1 2221 WKALGSVIVCSETVDIPLINDYQIEDINDFEELERGIQDEEL
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Results of SIM with:

Sequence 1: NP_598406.1 (2282 residues)

Sequence 2: BAB00956.1 (1283 residues)

using the parameters:

Comparison matrix: BLOSUM62

Number of alignments computed: 20

Gap open penalty: 12

Gap extension penalty: 4



Evaluate the significance of this protein sequence similarity score using PRSS at EMBOSS-CH

55.3% identity in 2264 residues overlap; Score: 11637.0; Gap frequency: 0.1%

```

NP_598406.1      1  MRASGETLLPEVHLNPTVYHKLIIYLLGHFPHDLGLSETSPFHNNNDQIAREESNLA
BAB00956.1      1  MRASGETLLPEVHLNPTVYHKLIIYLLGHFPHDLGLSETSPFHNNNDQIAREESNLA
      * * * * *
NP_598406.1     61  SELGVAKSSLIKEVPAPRATKWRSHAVLINPSCIPEFLYKFLPESKLIQVREQNYKLINAS
BAB00956.1     61  SELGVAKSSLIKEVPAPRATKWRSHAVLINPSCIPEFLYKFLPESKLIQVREQNYKLINAS
      * * * * *
NP_598406.1    121  CNTTSDSIDRCMELISIKLTGFNNLFPSRSPTAGAGKNSKITLNDIQSINSENYKQPVVS
BAB00956.1    121  CNTTSDSIDRCMELISIKLTGFNNLFPSRSPTAGAGKNSKITLNDIQSINSENYKQPVVS
      * * * * *
NP_598406.1    181  LNLITIKYQMFQLDNIHQSSQPTOLVHIVDTRACSLIVITPELVIQFELNSVIMYTFPEMT
BAB00956.1    181  LNLITIKYQMFQLDNIHQSSQPTOLVHIVDTRACSLIVITPELVIQFELNSVIMYTFPEMT
      * * * * *
NP_598406.1    241  LNVSTMPFGSMNVYALCTISHTLSPLGPRIDPLPSIVDELAQLQDPTVYKVIASFLESLEY
BAB00956.1    241  LNVSTMPFGSMNVYALCTISHTLSPLGPRIDPLPSIVDELAQLQDPTVYKVIASFLESLEY
      * * * * *
  
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NP-598406. 391 GCGCLQKDFVVELASPPHSFITERIDILIGSYALDYDESITWVTQLDIFENLSPOLLE
 EAB00056.1 391 GCLQKQDFVVELASPPHSFITERIDILIGSYALDYDESITWVTQLDIFENLSPOLLE
 NP-598406. 361 MLCCLMRANSHETLTQAQV-GKVRSGMAGULLDFTTINKTIAFPFTILINQYRKKKGMW
 EAB00056.1 361 MLCCLMRANSHETLTQAQAGNVPBMCAGKLLDFTTINKTIAFPFTILINQYRKKKGMW
 NP-598406. 420 FPLILFPMASKEELINFGHONASISYEVTLAGHWEELSTEPFKCFDPFGSELSIFMKDKA
 EAB00056.1 421 FPLILFPMASKEELINFGHONASISYEVTLAGHWEELSTEPFKCFDPFGSELSIFMKDKA
 NP-598406. 460 TSAFSDWMSVFRSLIDQRHQBHEIPDPFNRLLLEPLEDDSPDVAELRYVTOTY
 EAB00056.1 461 TSAFSDWMSVFRKLLKQPHQRHIMPDPFNRLLLEPLEDDSPDVAELRYVTOTY
 NP-598406. 540 LQDITFCASYLKEKEIKPLARIPAKLTMRHESQVIRBAILANAGTILAKNGVVLNQL
 EAB00056.1 541 LQDITFCASYLKEKEIKPDRIFAKLTMRHESQVIRBAILANAGTILAKNGVVLNQL
 NP-598406. 600 SUTBSLTHNQKSTIDRKAERYTDMISQSGPHTIKTUSPMKNGPTACSYLTDPGDTPE
 EAB00056.1 601 SUTBSLTHNQKSTIDRKAERYTDMISQSGPHTIKTUSPMKNGPTACSYLTDPGDTPE
 NP-598406. 660 LSACFHTTOLAKYCLQWEXQTIHPARTLNRMVGVPLFVEMILMLIASHLVGDDFPRP
 EAB00056.1 661 LSACFHTTOLAKYCLQWEXQTIHPARTLNRMVGVPLFVEMILMLIASHLVGDDFPRP
 NP-598406. 720 AATDAFDLQKVLNODIFIVE-RSGLEGLQKNWIMISTSVILSGRANCRVMSVQGLN
 EAB00056.1 721 AATDAFDLQKVLNODIFIVGPMHIEGLQKNWIMISTSVILSGRANCRVMSVQGLN
 NP-598406. 770 QALAVITRVFESLESIOKKELAYASKLFPBELRANNYGLGHOKAQSTISSTFFPIYK
 EAB00056.1 771 QALAVITRVFESLESIOKKELAYASKLFPBELRANNYGLGHOKAQSTISSTFFPIYK
 NP-598406. 839 RVFYQGRILUTQALNKAELCLTAGVLGECTQASCSMSATTMLNTEGVSKYTCUKLNTY
 EAB00056.1 841 RVFYQGRILUTQALNKAELCLTAGVLGECTQASCSMSATTMLNTEGVSKYTCUKLNTY
 NP-598406. 899 QSTHQLNYDLIFPQYSIFGHTISSEIFLQNPRLYSRTVLLPSQLGLIYVLACSLNENIG
 EAB00056.1 901 QSTHQLNYDLIFPQYSIFGHTISSEIFLQNPRLYSRTVLLPSQLGLIYVLACSLNENIG
 NP-598406. 959 DPLQTVADLKLPLIKGALSWSILYNLAKRPKGSWATLAADPSYLNQBYLPPFTILK
 EAB00056.1 961 DPLQTVADLKLPLIKGALSWSILYNLAKRPKGSWATLAADPSYLNQBYLPPFTILK
 NP-598406. 1019 RHTQNTLARIENPMLFOVFTMAKEENLLANPLLEDIVLPVAVHIIIGGSGIOMKKQ
 EAB00056.1 1021 RHTQNTLARIENPMLFOVFTMAKEENLLANPLLEDIVLPVAVHIIIGGSGIOMKKQ
 NP-598406. 1070 IQGFFYTITWIMASFPRIEMLOTKRTISVLENTNYLSYNYVILNPLFTIQLVYTTQ

BAR00055.1 1081 IQCFEFTTETIMPRSPFIFPLSTYKTLSEVIEYNINYLEYRIEVLAPLPIFNYLWFTIQ
 * * * * *

NP-598406. 1139 TCSIDISRLKRLSWSGLLNGKTLLEGLETFDFITFVFNSTLIVTODCLFCQSGDDCKTFW
 BAR00055.1 1141 TCSIDISRLKRLSWSGLLNGKTLLEGLETFDFITFVFNSTLIVTODCLFCQSGDDCKTFW
 * * * * *

NP-598406. 1199 FLPMGIIIDONFETNPPFIPVPIOSRTTEHRVAMAYIYGATHSLEKALRGASVYIMAFQ
 BAR00055.1 1201 FLPMGIIIDONFETNPPFIPVPIOSRTTEHRVAMAYIYGATHSLEKALRGASVYIMAFQ
 * * * * *

NP-598406. 1259 DTVVNMNMDLIEETVMIISLEQLQTLTLPTEBMITHELDKATTLKFTFASSTAFSSY
 BAR00055.1 1261 DTVVNMNMDLIEETVMIISLEQLQTLTLPTEBMITHELDKATTLKFTFASSTAFSSY
 * * * * *

NP-598406. 1319 TMSNQQLYLEIDGRVVDNIIYQGLMITGLGIIETYNHFFIARTSTQBITLMLATSSOC
 BAR00055.1 1321 TMSNQQLYLEIDGRVVDNIIYQGLMITGLGIIETYNHFFIARTSTQBITLMLATSSOC
 * * * * *

NP-598406. 1379 VRSVDSCLINESDGRVPIITVPIYHTFVYDPLADYSIAHLYLSYQAKKIGSTYYSLT
 BAR00055.1 1381 VRSVDSCLINESDGRVPIITVPIYHTFVYDPLADYSIAHLYLSYQAKKIGSTYYSLT
 * * * * *

NP-598406. 1439 DQIFLLGLTAKQMINSLIGLDETVEIVHDAVILSDYTRNWIISCSYTKIDLVPKIMRN
 BAR00055.1 1441 DQIFLLGLTAKQMINSLIGLDETVEIVHDAVILSDYTRNWIISCSYTKIDLVPKIMRN
 * * * * *

NP-598406. 1499 FLLELAFQNYLRISSWTNIFDYTYMILKRIPTGALNHTLATISHPFLRFANMLLITP
 BAR00055.1 1501 FLLELAFQNYLRISSWTNIFDYTYMILKRIPTGALNHTLATISHPFLRFANMLLITP
 * * * * *

NP-598406. 1559 IHAFYLASLDYFKLSIDAIGWYKQVIALGSLHGILKLLILSBDGSMISDRAHMLTKRL
 BAR00055.1 1561 IHAFYLASLDYFKLSIDAIGWYKQVIALGSLHGILKLLILSBDGSMISDRAHMLTKRL
 * * * * *

NP-598406. 1619 LLLALYKGNITTFKIKOMPPREKCLVITEYLAMCYQHTMLDPLQKLYLYHITWFLYA
 BAR00055.1 1621 LLLALYKGNITTFKIKOMPPREKCLVITEYLAMCYQHTMLDPLQKLYLYHITWFLYA
 * * * * *

NP-598406. 1679 FFSNNHYLTKILNQIRSDSGQYIITSYYSFQELTDITILMSTLXPHVNSHNSKVR
 BAR00055.1 1681 FFSNNHYLTKILNQIRSDSGQYIITSYYSFQELTDITILMSTLXPHVNSHNSKVR
 * * * * *

NP-598406. 1739 FIFDFLPPHESLEMYPLFVDNDGQALSTLIPGPPSHVLRFLGVSTAWYKHSYCKY
 BAR00055.1 1741 FIFDFLPPHESLEMYPLFVDNDGQALSTLIPGPPSHVLRFLGVSTAWYKHSYCKY
 * * * * *

NP-598406. 1799 LETQKIQTDHLYLBSGSMWSLELLELFPDITVYNLSFSGENFPQRXYLFTQPVQ
 BAR00055.1 1801 LETQKIQTDHLYLBSGSMWSLELLELFPDITVYNLSFSGENFPQRXYLFTQPVQ
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NP-598406. 1859 SVFVLMQADLAFTDLKIDFVFLHQRKGAVTGLSTHNAVAFIHKVGSBEKASLVHLE
 BAR00055.1 1861 SVFVLMQADLAFTDLKIDFVFLHQRKGAVTGLSTHNAVAFIHKVGSBEKASLVHLE

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NP-598406. 1915 GTANINQQTLGRSQINGLIATTVLKPGGILYKTSWLPSPFQGLAQLMCTFFKINHLI
BAR00056.1 1920 STANIRGQLTGRSQKHSLEIATTVLKEGGILYKTSWLPSPFQGLAQLMCTFFKINHLI
*****
NP-598406. 1979 ESSYSQPSSEHGVYLCRLAADPRTIGFSALLVTAITLNRGPTTIGPDVVCSTWQHLEN
BAR00056.1 1980 ESSYSQPSSEHGVYLCRLAADPRTIGFSALLVTAITLNRGPTTIGPDVVCSTWQHLEN
*****
NP-598406. 2039 WGRVGVIVIEILLOGLATNFFAGHNLILKCOOTPSRRKYLEIDQLASFGVLQDALVILIT
BAR00056.1 2040 YSRVGVKVIDEILLOGLATNFFAGHNLILKCOOTPSRRKYLEIDQLASFGVLQDALVILIT
*****
NP-598406. 2099 IHLKELLEVQSSHTEDYTSLLPTFYNIGAGKQRTYIKLILERSLMTVTVMVLVLPSSYR
BAR00056.1 2100 IHLKELLEVQSSHTEDYTSLLPTFYNIGAGKQRTYIKLILERSLMTVTVMVLVLPSSYR
*****
NP-598406. 2159 DSVFQDLRLQSFRLMSILSEQTFLAKTETTKYTLIDQHTETTYLSTFFNSHGVLMKRPYQK
BAR00056.1 2160 DSVFQDLRLQSFRLMSILSEQTFLAKTETTKYTLIDQHTETTYLSTFFNSHGVLMKRPYQK
*****
NP-598406. 2219 QJWKALGSVLYCSETVDIPLIKDIQIKYINDFPDLKRGIDQREL
BAR00056.1 2220 QJWKALGSVLYCSETVDIPLIKDIQIKYINDFPDLKRGIDQREL
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Supplemental Response to Examiner's Request

Applicants note that this response is being made at the request of the examiner; therefore, no reduction in any patent term extension due to Applicants should occur based on this response (see 37 CFR 1.704(c)(8)).

Date: June 7, 2010

/Sean C. Brock/

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